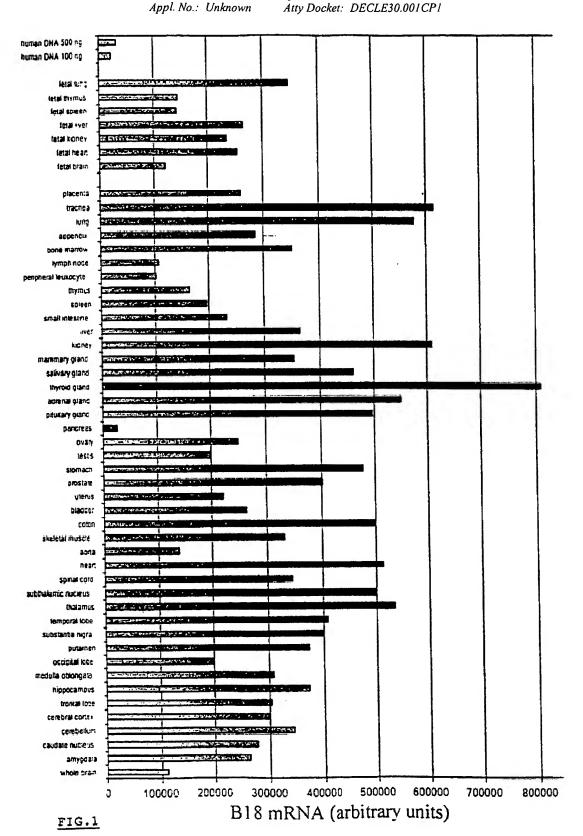
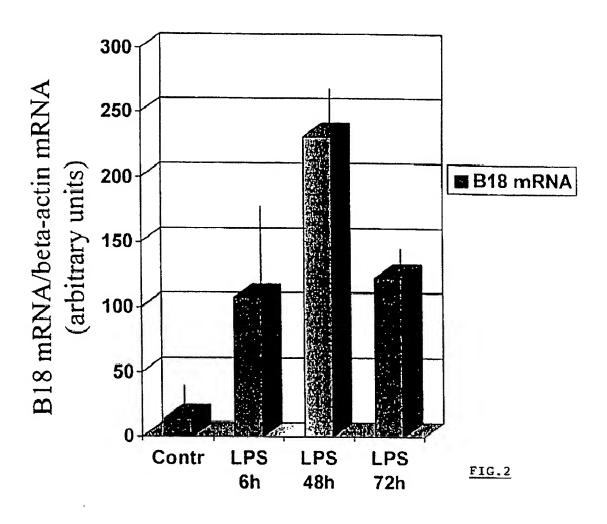
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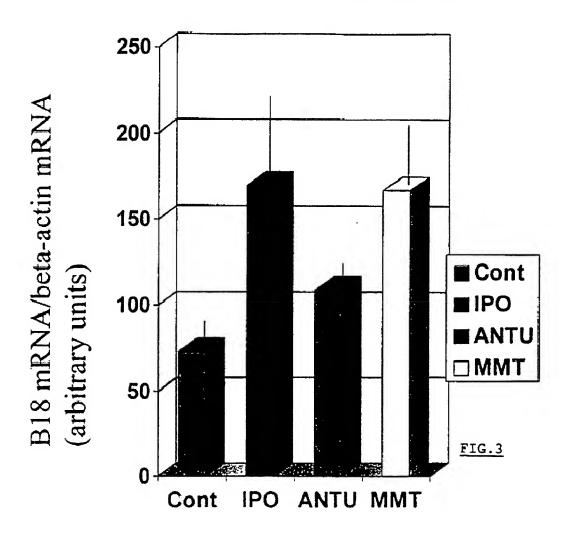
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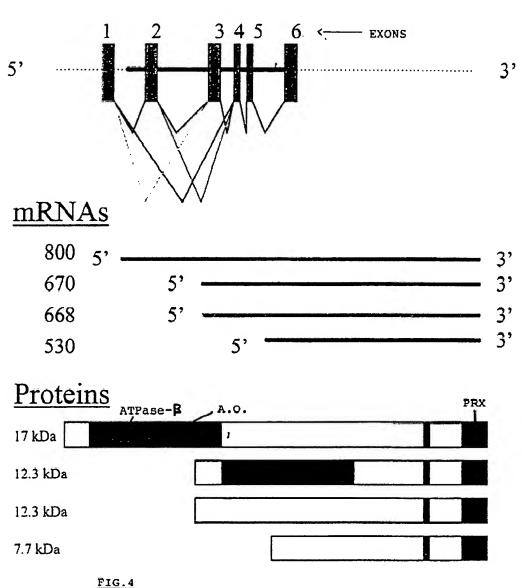


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Gene (chromosome 11q12-13)



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CLUSTAL V alig	nment of	human and	i rat	B18	amino	acid	sequences	(Identity:
90%, Homology:	97.5%):							

B18hum B18rat	MAPIKVGDAIPAVEVPEGEPGNKVNLAELFKGKKGVLFGVPGAFTPGCSK MAPIKVGDTIPSVEVFEGEPGKKVNLAELFKDKKGVLFGVPGAFTPGCSK	SEQIDNO1
B18hum B18rat	THLPGFVEQAEALKAKGVQVVACL5VNDAFVTGEWGRAHKAEGKVRLLAD THLPGFVEQAGALKAKGAQVVACL5VNDVFVTAEWGRAHQAEGKVQLLAD	FIG.5a
Bl8hum Bl8rat	PTGAFGKETDLLLDDSLVSIFGNRRLKRFSMVVQDGIVKALNVEPDGTGL PTGAFGKETDLLLDDSLVSLFGNRRLKRFSMVIDKGVVKALNVEPDGTGL	
B18hum B18rat	TCSLAPNIISQL TCSLAPNILSQL	

CLUSTAL V alignment of human and mouse B18 amino acid sequences (Identity: 91%, Homology: 96%):

Bl8mouse	MAPIKVGDAIPAVEVFEGEPGNKVNLAELFKGKKGVLFGVPGAFTPGCSK MAPIKVGDAIPSVEVFEGEPGKKVNLAELFKGKKGVLFGVPGAFTPGCSK
B18hum B18mouse	THLPGFVEQAEALKAKGVQVVACLSVNDAFVTGEWGRAHKAEGKVRLLAD THLPGFVEQAGALKAKGAQVVACLSVNDVFVIEEWGRAHQAEGKVRLLAD
B18hum	PTGAFGKETDLLLDDSLVSIFGNRRLKRFSMVVQDGIVKALNVEPDGTGL
B18mouse	PTGAFGKATDLLLDDSLVSLFGNRRLKRFSMVIDNGIVKALNVEPDGTGL
B18hum	TCSLAPNIISQL
B18mouse	TCSLAPNILSQL

CLUSTAL V alignment of human and rat cDNA sequences (identity: 612/780, 78.5%):

B18hum B18rat	GCCAGGAGGCGGAGTGGAAGTGGCCGTGGGGCGGGTATGGGACTAGCTGG
B18hum B18rat	CGTGTGCGCCCTGAGACGCTCAGCGGGCTATATACTCGTCGGTGGGGCCG
Bl8hum Bl8rat	GCGGTCAGTCTGCGGCAGCGGCAGCAAGACGGTGCAGTGAAGGAGAGTGG

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	A I M II I A D I D D C CO COLOR
FIG.5b	Appl. No.: Unknown Atty Docket: DECLE30.001CP1
B18hum	GCGTCTGGCGGGGTCCGCAGTTTCAGCAGAGCCGCTGCAGCCATGGCCCC
B18rat	GAGTGTGGTGGGGCCCGCAGCTTCAGCAGTGCCGCGGTGACTATGGCCCC
B18hum	AATCAAGGTGGGAGATGCCATCCCAGCAGTGGAGGTGTTTGAAGGGGAGC
B18rat	GATCAAGGTGGGAGACACCATTCCCTCAGTGGAGGTATTTGAAGGGGAAC
B18hum	CAGGGAACAAGGTGAACCTGGCAGAGCTGTTCAAGGGCAAGAAGGGTGTG
B18rat	CTGGAAAGAAGGTGAACTTGGCAGAGCTGTTCAAGGACAAGAAAGGTGTT
B18hum	CTGTTTGGAGTTCCTGGGGCCTTCACCCCTGGATGTTCCAAGACACACCT
B18rat	TTGTTTGGAGTCCCTGGGGCATTTACACCTGGCTGTTCCAAGACCCATCT
B10hum	GCCAGGGTTTGTGGAGCAGGCTGAGGGCTCTGAAGGCCAAGGGAGTCCAGG
B10rat	GCCTGGGTTTGTGGAGCAAGCCGGAGCTCTGAAGGCCAAGGGAGCACAAG
B18hum	TGGTGGCCTGTCTGAGTGTTAATGATGCCTTTGTGACTGGCGAGTGGGGC
B18rat	TGGTGGCCTGTCTGAGTGTTAATGATGTCTTCGTGACTGCAGAGTGGGGT
B18hum	CGAGCCCACAAGGCGAAGGCAAGGTTCGGCTCCTGGCTGATCCCACTGG
B18rat	CGAGCCCACCAGGCAGAGGCAAGGTTCAGCTCCTGGCTGACCCCACTGG
Blőhum	GGCCTTTGGGAAGGAGACAGACTTATTACTAGATGATTCGCTGGTGTCCA
Blőrat	AGCTTTTGGAAAGGAGACAGATTTACTACTAGATGATTCTTTGGTGTCTC
B18hum	TCTTTGGGAATCGACGTCTCAAGAGGTTCTCCATGGTGGTACAGGATGGC
B18rat	TCTTTGGGAATCGTCGGCTAAAAAGGTTCTCCATGGTGATAGACAAGGGC
Bl8hum	ATAGTGAAGGCCCTGAATGTGGAACCAGATGGCACAGGCCTCACCTGCAG
Bl8rat	GTAGTAAAGGCACTGAACGTGGAGCCGGATGGCACAGGCCTCACCTGCAG
B18hum	CCTGGCACCCAATATCATCTCACAGCTCTGAGGCCCTGGGCCAGATTACT
B18rat	CCTGGCCCCCAACATCCTCTCACAACTCTGAGGCCCTGA-CCAGAATG
B18hum B18rat	TCCTCCACCCCTCCCTATCTCACCTGCCCAGCCCTGTGCTGG-GGCCCTG TCCTCTGACTCTCCC-ATCTCCTCCACCCAGCTCTGGGCCAAAGGCCCAG
B18hum B18rat	CATTGGCCAGATTTCTGC TACCTCCTTACCTGAGGGCCACTGGAATGGAA
B10hum B10rat	AATAAACACTTGTGGTTTGCGGAAAAAAAAAAAAAAAAA

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CLUSTAL V alignment of human and mouse cDNA sequences (Identity: 552/675, 81.8%):

01.05):	FIG.5c
B18hum B18mouse	GCCAGGAGGCGGAGTGGAAGTGGCCGTGGGGGGGGGTATGGGACTAGCTGG
Bl8hum Bl8mouse	CGTGTGCGCCCTGAGACGCTCAGCGGGGCTATATACTCGTCGGTGGGGCCG
Bl8hum Bl8mouse	GCGGTCAGTCTGCGGCAGCGGCAGCAAGACGGTGCAGTGAAAGAAGCAGGTTGG
B18hum	GCGTCTGGCGGGGTCCGCAGTTTCAGCAGAGCCGCTGCAGCCATGGCCCC
B18mouse	GAGTGTGGCGGAGCCCGCAGCTTCAGCAGCTCCGCGGTGACCATGGCCCC
B18hum B18mouse	AATCAAGGTGGGAGATGCCATCCCAGCAGTGGAGGTGTTTGAAGGGGAACGATCAAGGTGGAGATGCCATTCCCTCAGTGGAGGTATTTGAAGGGGAAC
818hum	CAGGGAACAAGGTGAACCTGGCAGAGCTGTTCAAGGGCAAGAAGGGTGTG
B18mouse	CGGGAAAGAAGGTGAACTTGGCAGAGCTGTTCAAGGGCAAGAAAGGTGTT
B18hum	CTGTTTGGAGTTCCTGGGGCCTTCACCCCTGGATGTTCCAAGACACACCT
B18mouse	TTGTTTGGAGTCCCTGGGGCATTTACACCTGGCTGTTCTAAGACCCACCT
Bl8hum	GCCAGGGTTTGTGGAGCAGGCTGAGGCTCTGAAGGCCAAGGGAGCCCAGG
Bl8mouse	GCCTGGGTTTGTGGAGCAAGCTGGAGCTCTGAAGGCTAAGGGAGCGCAGG
B18hum	TGGTGGCCTGTCTGAGTGTTAATGATGCCTTTGTGACTGGCGAGTGGGGC
B18mouse	TGGTGGCCTGTCTGAGCGTTAATGACGTCTTTGTGATTGAAGAGTGGGGT
B18hum	CGAGCCCACAAGGCGGAAGGCAAGGTTCGGCTCCTGGCTGATCCCACTGG
B18mouse	CGAGCCCACCAGGCAGAAGGCAAGGTTCGGCTCCTGGCTGACCCCACTGG
Bl8hum	GGCCTTTGGGAAGGAGACAGACTTATTACTAGATGATTCGCTGGTGTCCA
Bl8mouse	AGCCTTTGGGAAGGCGACAGACTTATTATTGGATGATTCTTTGGTGTCTC
Bl8hum	TCTTTGGGAATCGACGTCTCAAGAGGTTCTCCATGGTGGTACAGGATGGC
Bl8mouse	TCTTTGGGAATCGTCGGCTGAAAAGGTTCTCCATGGTGATAGACAACGGC
B18hum	ATAGTGAAGGCCCTGAATGTGGAACCAGATGGCACAGGCCTCACCTGCAG
B18mouse	ATAGTGAAGGCACTGAACGTGGAGCCAGATGGCACAGGCCTCACCTGCAG
Bl8hum	CCTGGCACCCAATATCATCTCACAGCTCTGAGGCCCTGGGCCAGATTACT
Bl8mouse	CCTGGCCCCCAACATCCTCTCCCAACTCTGAGGCCCTGG-CCAGATG
B18hum B18mouse	TCCTCCACCCCTATCTCACCTGCCCAGCCCTGTGCTGGGGCCCTGC TCCTCTGACTCTCCC-ATCTCTCCCACCCGGCTCTAGGCC
B18hum	AATTGGAATGTTGGCCAGATTTCTGCAATAAACACTTGTGGTTTGCGGAA
B18mouse	AAAAGGCTCGGTACCTCCTTACTGGGAGC-CACGT

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Fig. 6.

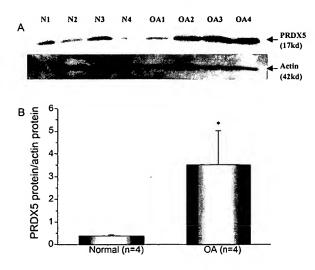
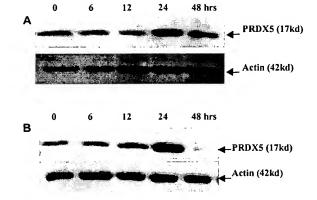


Fig. 7.



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Fig. 8.

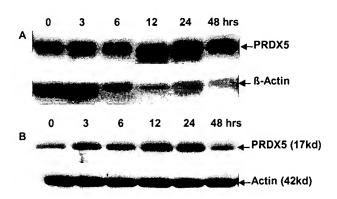
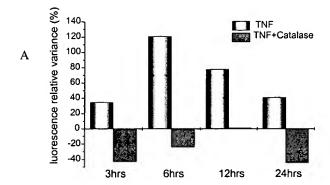
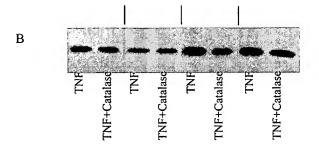
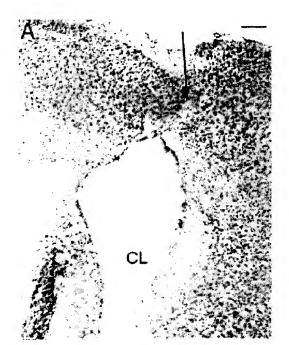


Fig. 9.

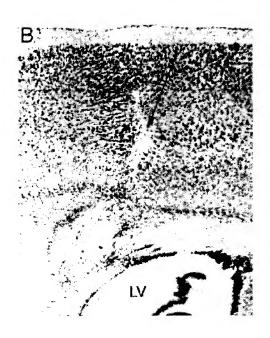




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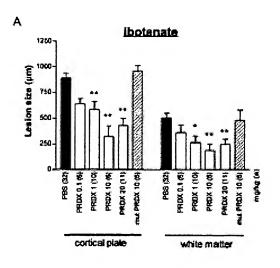


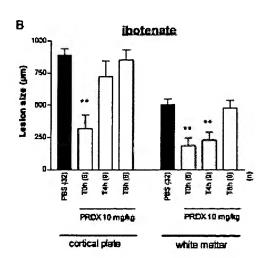


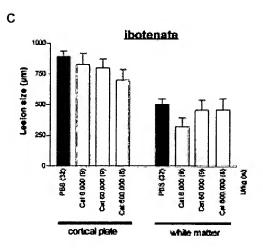


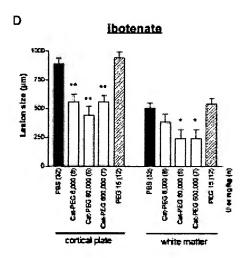
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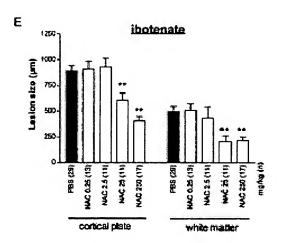
Fig. 11











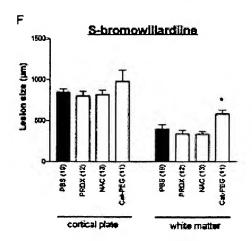
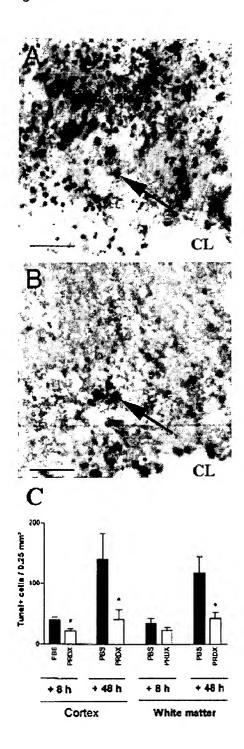
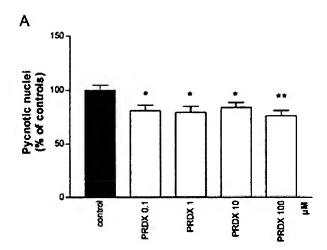
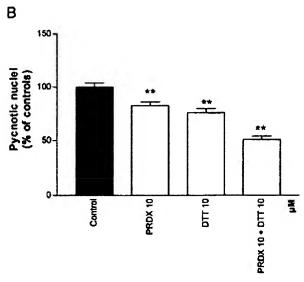


Fig. 12







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Fig. 14

